## PROPOSED CLAIMS FOR U.S. PATENT APPLN. NO. 10/539,834 Your Ref: CH:YN, PC/S-84-5US; Our Ref: 159-89

Claims 1-12 (canceled)

- 13. (currently amended) A method for transferring N-acetyl-D-glucosamine from a donor substrate to an acceptor substrate through  $\beta$ 1,3-linkage, wherein " $\beta$ " represents an anomer assuming a cis configuration, of anomers of glycosidic linkage at position 1 of the sugar ring, the method comprising reacting the donor substrate and the acceptor substrate in the presence of [[with]] a  $\beta$ 1,3-N-acetyl-D-glucosaminyltransferase protein, wherein the protein comprises the amino acid sequence of SEQ ID NO: 2 or SEQ ID NO: 16.
- 14. (currently amended) An isolated  $\beta$ 1,3-N-acetyl-D-glucosaminyltransferase protein comprising the amino acid sequence of SEQ ID NO: 2 or SEQ ID NO: 16, wherein the protein has the following properties (a) to (c):
- (a) acceptor substrate specificity: the protein transfers GlcNAc from UDP-GlcNAc to an acceptor substrate having an oligosaccharide residue in a quadruple-stranded form at the nonreducing end of an N-linked oligosaccharide through a β1,3 glycosidic linkage and synthesizes an oligosaccharide wherein "GlcNAc" represents an N-acetyl-D-glucosamine residue, and "β" represents an anomer assuming a cis configuration, of anomers of glycosidic linkage at position 1 of the sugar ring;
- (b) reaction pH: the protein has a high activity at or around neutral; and
- (c) divalent ion requirement: the activity is enhanced in the presence of at least Mn<sup>2+</sup> or Co<sup>2+</sup>.

Claims 15-30 (canceled)

31. (previously presented) The method according to Claim 13, wherein the protein comprises the amino acid sequence of SEQ ID NO: 2.

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- 32. (previously presented) The method according to Claim 13, wherein the protein comprises the amino acid sequence of SEQ ID NO: 16.
- 33. (previously presented) The protein of Claim 14 comprising the amino acid sequence of SEQ ID NO: 2.
- 34. (previously presented) The protein of Claim 14 comprising the amino acid sequence of SEQ ID NO: 16.

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